

Amendments to the specification:

On page 1, please insert the following paragraph following the title of the invention:

The present application is a Divisional Application of USSN 09/257,069, filed February 24, 1999, which issued as US Patent 6,348,580 on February 19, 2002, which in turn claimed the priority benefit of Japanese Patent Application No. 10-252921, filed September 7, 1998.

On page 33, please amend the paragraph appearing at lines 10-21 as follows:

The amino acid sequence for the thus obtained gene encodes was deduced, and hypervariable regions CDR were identified by the Chothia Numbering Scheme (<http://www.biochem.ucl.ac.uk/~martin/abs/GeneralInfo.html#kabatnum>, Al-Lazikani et al., J. Molec. Biol., 273, 927-948, 1997). The amino acid sequences constructing the thus identified variable regions of the heavy chains, light chains, and CDRs are shown in Fig. 8. The amino acid residues marked with the arrow in Fig. 8 hardly appear at these positions in conventional antibodies, according to the antibody database. The amino acid residues located in the CDRs thus seem to participate in the specificity. In order to obtain mutants of the amino acid sequences, the sequence containing the amino acid residues should be selected.